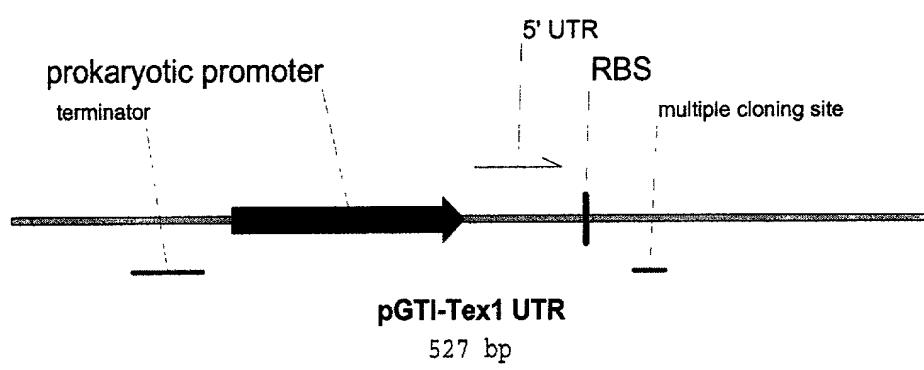


**Fig. 1**



**Fig. 2**

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GGCCGGGGCCCCGCCCCCTTGGCGGGGCCTCCCCCAAGGAGGGCCG

**Fig. 3**

GCATGCTTATCTCGAGACTGGCAGTTCAATAGAGATATTGTATGCCTGCAG

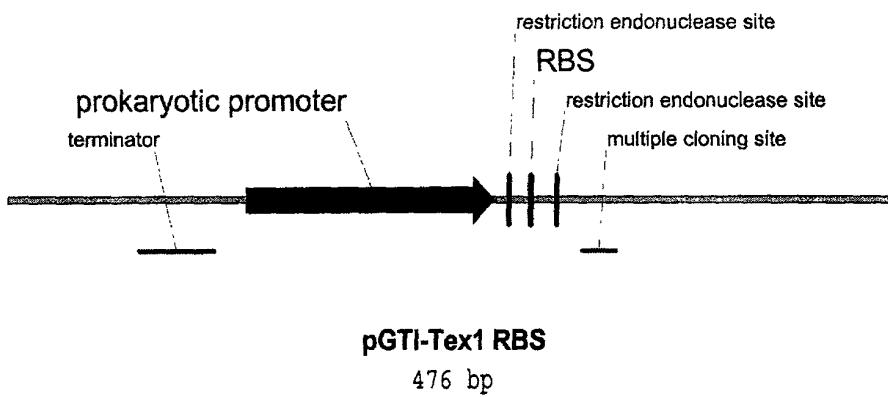
**Fig. 4**

AAAGGGA

**Fig. 5**

GAAGGAGG

**Fig. 6**



**Fig. 7**

T006405 \* 02000

GGAGGAGGCC AAGAAGCTTT TGGAGGGAA GCCCGTCTAC ATGTACCCCCA CGTCATTGA	60
GGCGGCCAAG GCCATCGTGG CCATGGTGGG AGGTGCGGCG TGATCCTGGT GAACCGCGAG	120
ACCCGCGTCC TGGTCCAGGG CATCACCGGC CGGGAGGGGC AGTTCCACAC CAAGCAGATG	180
CTGGACTACG GCACCAAGAT CGTCGCCGG GTCACCCCGG GCAAAGGGGG AACGGAGGTC	240
CTAGGGGTCC CCGTCTACGA CACGGTGAAG GAGGCGGTGG CGCACCACGA GGTGGACGCC	300
TCCATCATCT TCGTGCCCAGC CCCGGCCGCG GCGGACGCCG CCCTGGAAGC GGCCCACGCC	360
GGGATCCCCC TCATCGTCCT CATCACCGAG GGCACTCCCCA CCCTGGACAT GGTGCGGGCG	420
GTGGAGGAGA TCAAGGCCCT GGGAAAGCCGC CTCATCGGGG GGAAGTGCCT GGGGATCATC	480
AGCGCCGAGG AGACCAAGAT CGGGATCATG CCCGGCCACG TCTTCAAGCG GGGCCGGGTG	540
GGGATCATCA GCCGCTCCGG CACCCCTCACC TACGAGGCCG CAGCCGCCCT TTCCCCAGGCG	600
GGGCTCGGCA CCACCAACAC GGTGGGGATC GGGGGCGACC CCGTCATCGG CACCACCTTC	660
AAGGACCTCC TCCCCCTCTT CAACGAGGAC CCGGAGACGG AGGCCGTGGT CCTCATCGGG	720
GAGATCGGCG GCTCCGACGA GGAGGAGGCG GCGGCTTGGG TGAAGGACCA CA	772

**Fig. 8**